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**ERROR REPORT**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/562,601  
Source: IFWO  
Date Processed by STIC: 3/3/07

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FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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**<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>**

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1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
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Revised 01/10/06



IFWO

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION:** US/10/562,601

**DATE:** 03/05/2007  
**TIME:** 13:44:39

**Input Set :** N:\efs\03\_05\_07\10562601\_efs\sequencelisting.txt  
**Output Set:** N:\CRF4\03052007\J562601.raw

3 <110> APPLICANT: Hengst den, Christiaan D.  
 4 Gajic, Olivera  
 5 Kuipers, Oscar P.  
 6 Kok, Jan  
 7 Sikkema, Jan  
 8 Geurts, Johannes M.W.  
 9 Nauta, Arjen

11 <120> TITLE OF INVENTION: Methods and means for regulating gene expression  
 13 <130> FILE REFERENCE: P63590US00  
 15 <140> CURRENT APPLICATION NUMBER: US 10/562,601  
 16 <141> CURRENT FILING DATE: 2005-12-28  
 18 <150> PRIOR APPLICATION NUMBER: PCT/NL2004/000474  
 19 <151> PRIOR FILING DATE: 2004-07-02  
 21 <150> PRIOR APPLICATION NUMBER: EP 03077074.7  
 22 <151> PRIOR FILING DATE: 2003-07-02  
 24 <160> NUMBER OF SEQ ID NOS: 234  
 27 <170> SOFTWARE: PatentIn Ver. 3.3

*Does Not Comply  
 Corrected Diskette Needed*

*see p 2,4*

#### ERRORED SEQUENCES

2299 <210> SEQ ID NO: 179  
 2300 <211> LENGTH: 26  
 2301 <212> TYPE: DNA  
 2302 <213> ORGANISM: Artificial Sequence  
 2304 <220> FEATURE:  
 2305 <223> OTHER INFORMATION: presence of the putative CodY box in arcA  
 E--> 2307 <400> SEQUENCE: 179 *see p 2*  
 2308 atatttttg aaaaattttt taaaaaa

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<210> 179  
<211> 26  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> presence of the putative CodY box in arcA

e <400> 179  
atatttttg aaaaattttt taaaaaa

26

*delete extra opening bracket*

**VERIFICATION SUMMARY**

PATENT APPLICATION: US/10/562,601

DATE: 03/05/2007

TIME: 13:44:40

Input Set : N:\efs\03\_05\_07\10562601\_efs\sequencelisting.txt  
Output Set: N:\CRF4\03052007\J562601.raw

L:2307 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:179 differs:178  
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L:2744 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2737  
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2742  
L:2750 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:214,Line#:2747  
L:2914 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:227 after pos.:0  
L:2931 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:228 after pos.:0  
L:2960 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:230 after pos.:0

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<210> 15  
<211> 8  
<212> PRT  
<213> Artificial Sequence

<220>  
<223> consensus sequence GTP binding motif in CodY homologs

<220>  
<221> Variant  
<222> (2)..(2)  
<223> Arg

if the amino acid at this location can represent  
another amino acid, use "Xaa" instead, and  
explain in <220>-<223> section

<220>  
<221> Variant  
<222> (3)..(3) same  
<223> Met, Gly, Ile, Lys, Gln

<220>  
<221> Variant  
<222> (4)..(4) same  
<223> Gly

<220>  
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<222> (5)..(5) same  
<223> Thr

<220>  
<221> Variant  
<222> (7)..(7) same  
<223> Ser

<400> 15  
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1 5